

Figure 1 Gregg, R.K., et. al.

91A₃H

_____ D _____

SEQ ID NO. 5 TYR PHE CYS ALA ARG SER TYR TYR SER GLY ASP MET TYR CYS
 SEQ ID NO. 7 TAT TTC TGT GCA AGA TCG TAT TAC TCT GGT GAT ATG TAC TGC

SEQ ID NO. 6 PHE ASP TYR TRP
 SEQ ID NO. 8 TTT GAC TAC TGG

91A₃H-INSβ

_____ INSβ _____

SEQ ID NO. 9 - - - - - HIS LEU VAL GLU ALA LEU - LEU VAL CYS GLY GLU ARG GLY - - - - -
 SEQ ID NO. 10 - - - - - AGC C-C CTA GTG -AG -CG C-T --- CT- GTT TGC GGT GAA AGA GGT - - - - -

91A₃H-HEL

_____ HEL _____

SEQ ID NO. 11 - - - - - ALA MET LYS ARG HIS GLY LEU ASP ASN TYR ARG GLY TYR SER LEU - - - - -
 SEQ ID NO. 12 - - - - - GCA ATG A-G CGC CAC -GG A-A G-- AA- TAT CGG GGA TAT AGC CTC - - - - -

